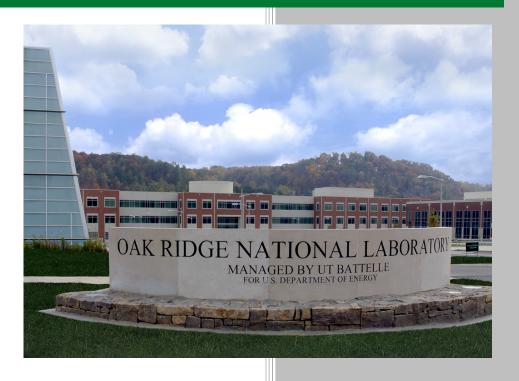
Acceleration of Thermochimica Calculations in Bison



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Computational Sciences & Engineering Division

Acceleration of Thermochimica Calculations in Bison

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ABSTRACT

The performance of the open-source thermochemical software library Thermochimica has been enhanced by developing a re-initialization algorithm to make use of data from previous calculations to reduce the number of convergence steps in the Gibbs energy minimization procedure. This algorithm has been tested in the context of stand-alone Thermochimica calculations, and speedups in the range of 2x-3x achieved for cases with chemistries resembling those of irradiated nuclear fuels. Routines to make use of this re-initialization procedure have been implemented in Bison, in which each node uses the results of the previous calculation at that node as the initial conditions for the following Thermochimica call. Examples based on calculating the diffusion of oxygen in UO₂ Light Water Reactor (LWR) fuel in 1D and 3D have been tested, with speedups up to 7.38x demonstrated.

1. INTRODUCTION

The open-source thermochemical library Thermochimica [7, 11] has previously been coupled with the Multi-physics Object-Oriented Simulation Environment (MOOSE) [3] Finite Element Method (FEM) app Bison [5] for the purpose of providing thermodynamic information to simulations of nuclear fuels, such as phase fractions, chemical potentials, concentrations of species, vacancies and interstitials. In the initial implementation, each call to Thermochimica by Bison was independent of all previous such calls, and each thermochemical calculation begins anew. However, in typical Bison simulations, the change in system parameters (i.e., chemical element concentrations, temperature, and pressure) between calls to Thermochimica is very small, and thus the calculation results are typically very similar. Here, we implement a method for initializing Thermochimica calculations by making use of the results from previous calculations, and demonstrate that considerable speed-up is achieved when Bison simulations make use of this new initialization method. This speedup is crucial for simulations of nuclear fuels, which require trillions of thermochemical calculations due to the large physical dimensions and long timescales involved in these applications. Further, nuclear fuel systems tend to be chemically complex (with 20 or more chemical elements often involved), which leads to each call to Thermochimica being highly computationally expensive.

In a stand-alone Thermochimica calculation, the leveling algorithm [2] is used to provide an initial estimation of the phase assemblage and the element potentials. The leveling method is a first approximation to equilibrium, which does not include mixing between species in a solution phase – all species and phases are temporarily treated as stoichiometric phases. Once this initial estimate of phase assemblage has been obtained, the Gibbs Energy Minimization (GEM) method [16] is used to determine the combination of phases and phase compositions that yields the lowest integral Gibbs energy. This is done by iteratively manipulating the quantities of species to minimize the integral Gibbs energy, adding to the assemblage those phases with a negative driving force, and removing phases from the assemblage that tend to form negative (i.e., non-physical) quantities.

In the new method implemented here, the first Thermochimica calculation at each point of interest in the Bison system (i.e., nodes or integration points) proceeds as above. However, immediately after each calculation, key chemical information provided by Thermochimica is stored by Bison on a mesh. Then, on following calls to Thermochimica, that information is reused to provide an enhanced estimation of the initial phase assemblage and species concentrations. This has two positive effects: the leveling solver may be bypassed, and (more significantly) the resulting assemblage and concentrations are typically much

closer to the equilibrium state than those provided by the leveling solver. When the GEM solver begins with a better initial estimate, it takes many fewer steps to converge, and the calculation time is greatly decreased. Of course, storage of this initialization data at each node or integration point results in increased total memory usage as well as additional memory operations; however, our benchmarks demonstrate that these costs are greatly outweighed by the reduced number of calculation cycles.

In this report, we will provide a brief overview of the standard Thermochimica calculation method (i.e. leveling procedure and GEM solver), describe the new initialization method and its implementation in Bison, and finally discuss the example problems used to benchmark the speedup provided by the improvements described here.

2. THERMOCHIMICA OVERVIEW

Thermochimica is a software library that determines the equilibrium combination of phases and compositions of those phases given input temperature, pressure, and concentrations of chemical elements. Thermochimica and the development of Thermochimica as well as its integration into the MOOSE/Bison framework have been well-documented in previous technical reports [7, 14, 13, 12], and journal articles [11, 8, 10, 9], as well as in the Thermochimica documentation (available on GitHub).

From its genesis, Thermochimica has been designed to perform efficiently when coupled to multi-physics codes. This is reflected in the two-step algorithm design consisting of a leveling procedure followed by GEM. The leveling solver efficiently provides an accurate initial estimate of the phase assemblage by ignoring the composition dependence of the chemical potential of all solution species. Quickly obtaining this estimate allows the GEM routine to begin much closer to the correct equilibrium phase assemblage and phase compositions than if it were initialized randomly; thus, a large number of unnecessary GEM iterations are reduced.

The approach of the GEM method is to optimize the system such that the residuals in mass balance equations and the minimization of total Gibbs energy are simultaneously minimized, subject to the constraints that the Gibbs Phase Rule is respected and conservation of mass [16, 11]. This leads to the Lagrangian function [15]:

$$L = \frac{G}{RT} - \sum_{j=1}^{C} \tilde{\Gamma}_j (b_j - b_j^m)$$
 (1)

where G is the total Gibbs energy, R is the ideal gas constant, T is the temperature, C is the number of components* in the system, $\tilde{\Gamma}_j$ is the dimensionless chemical potential of component j, b_j is the number of gram-atoms of component j, and m is the GEM iteration index. Each GEM solver iteration requires the calculation of the Hessian (second partial derivative matrix) of L, which forms an $N \times N$ system where $N = C + \Phi$, where Φ is the total number of stable phases. This elucidates why the GEM solver is the most expensive component of Thermochimica as well as why calculation expense scales rapidly with the number of system components in the system.

As an equilibrium thermodynamics code, Thermochimica explicitly does not consider chemical kinetics, but rather assumes instantaneous local equilibrium. While this assumption is not strictly accurate, it is not a poor approximation for nuclear fuel applications because in these systems the temperature is typically high

^{*}The system components are typically taken as the chemical elements, but can be interpreted as integer combinations of the chemical elements.

and chemical elements are well mixed due to the stochastic nature of nuclear fission. In multiphysics simulations involving Thermochimica and Bison, all kinetics are handled within Bison, and Thermochimica calculations consider only the instantaneous state of the system when called. However, the kinetics are driven by variables provided by Thermochimica, such as the chemical potentials of diffusing species. This approach follows previous work in coupling Thermochimica with Bison for calculations of oxygen diffusion in nuclear fuels [14].

3. RE-INITIALIZATION METHOD

While the leveling procedure has proven to be a major performance enhancer for Thermochimica, neglecting the compositional dependence of chemical potential leads to discrepancies in the estimated phase assemblage, compositions, and chemical potentials that are in many cases greater than the changes in those variables between two consecutive evaluations in time within a multi-physics framework. Even small improvements to the accuracy of the initial values of these variables can lead to a considerable reduction in the number of trial assemblages and compositions calculated within the GEM subroutine, which is by far the most computationally expensive section of Thermochimica. When the savings in GEM iterations (plus the relatively small cost of the leveling solver) outweigh the overhead of storing and loading chemical data, computational efficiency can be improved by re-initializing Thermochimica with data from the previous calculation rather than making use of the leveling solver.

To bypass the leveling solver in Thermochimica, the arrays iAssemblage, dMolesPhase, dElementPotential, dMolFraction, and dChemicalPotential must be stored in memory and supplied in a subsequent call. The physical meanings, types, and sizes of these variables are described in Table 1. Additionally, an integer array created just for restarts, iElementsUsed, must be created and stored. iElementsUsed lists which elements were present in any non-zero amount in the calculation for which data was saved. If this differs from the current collection of elements, the restart attempt is aborted. The suffix _Old is added to each parameter when they are saved, and the variables with this prefix are held in the module ModuleRestart. Additional flags to track the status of the restart procedure are also available. These include iRestartRequested, which the user must set to .TRUE. if restarting is desired, iRestartAvailable, which is set to .TRUE. after restart data is successfully loaded, and iRestartLoaded, which is set to .TRUE. after a restart is successfully loaded.

Table 1. Summary of variables required to re-initialize a Thermochimica calculation

Variable Name	Array Dimension	Description
nElements	1	An integer scalar representing the number of
		elements in the system.
nSpecies	1	An integer scalar representing the number of
		species in the system.
iAssemblage	1:nElements	An integer vector representing the indices of
		the phases predicted to coexist at equilibrium.
dMolesPhase	1:nElements	A double real vector representing the number
		of moles of coexisting phases at equilibrium
		(corresponds directly to iAssemblage).
dElementPotential	1:nElements	A double real vector representing the element
		potentials.
dMolFraction	1:nSpecies	A double real vector representing the mole
		fraction of each species in the system data-
		file.
dChemicalPotential	1:nSpecies	A double real vector representing the chemi-
		cal potentials of all species in the system.

4. BISON IMPLEMENTATION

The implementation of Thermochimica re-initialization in Bison builds upon previous efforts to couple Thermochimica calculations into the MOOSE framework [14, 13, 12]. In addition to the creation of new interfaces to accommodate re-initialization, the existing Thermochimica implementations were generalized and recast into the form of a UserObject during this work. This will enable Bison users to make use of Thermochimica for any future simulations with minimal software development effort. The base UserObject, as well as an example derived UserObject and corresponding AuxKernel used for the demonstration problems discussed in Section 5.2 are described below.

4.1 ThermochimicaNodalData (MOOSE UserObject)

This class provides the base interface with Thermochimica for Bison. The input parameters are a vector containing the amount of each element, scalars for pressure and temperature, and optionally a switch for turning off the re-initialization algorithm (this algorithm is used by default). On each execute (which occurs when specified by the user on input), this UserObject inputs the temperature, pressure, and element concentrations to Thermochimica as well as any available re-initialization data, calls Thermochimica, then stores re-initialization data for the next time-step at each node in the FEM mesh.

4.2 ThermochimicaNodalUO2X (MOOSE UserObject)

The ThermochimicaNodalUO2X UserObject is an example specialization of the ThermochimicaNodalData UserObject described previously, designed to perform calculations for the

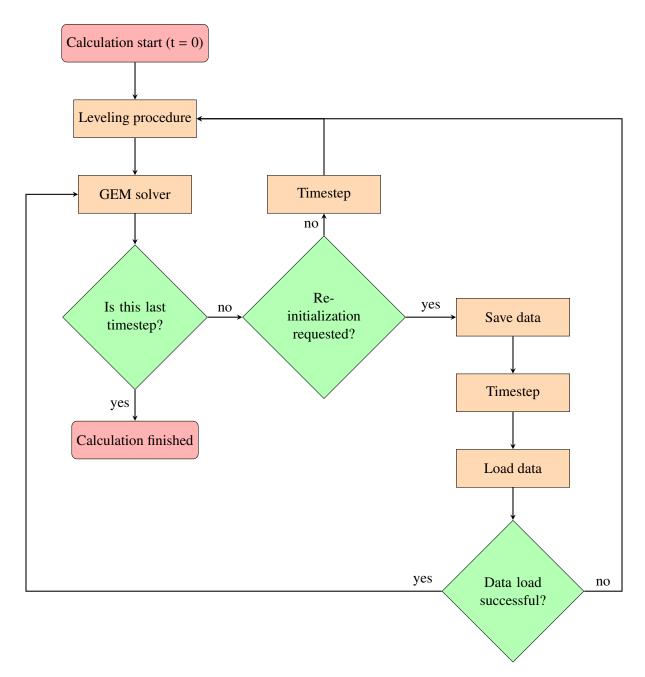


Figure 1. Flowchart outlining re-initialization procedure.

 $UO_{2\pm x}$ system. This UserObject makes use of the PresetElementMass function in the Thermochimica interface to overwrite the mass of oxygen from the elements AuxVariable used in the base ThermochimicaNodalData with the value in the oxygen variable. The Thermochimica calculation is performed by the base class, and then Thermochimica outputs of interest are extracted via various accessor functions provided by the Thermochimica interface. These are then saved in member variables and made available to other objects via get- accessor functions.

Future MOOSE/Bison objects that will interface with Thermochimica can be written following the example provided by this class. In particular, these should make use of the program structure used in execute() here:

- 1. preset masses for any elements for which this is necessary,
- 2. call base class execute(),
- 3. extract any required Thermochimica outputs and make them available to other objects.

4.3 UO2XChemistryTransportSimplified (MOOSE AuxKernel)

U02XChemistryTransportSimplified is an example of how results from the Thermochimica calculations performed by UserObjects that are specializations of ThermochimicaNodalData can be used to calculate values of AuxVariables. In this case, ThermochimicaNodalU02X is a UserObject that extracts data such as oxygen interstitial and vacancy site fractions, and chemical potential from a Thermochimica calculation. This AuxKernel copies that data to the relevant AuxVariables, which are later used to calculate the chemical potential and mobility of oxygen within $UO_{2\pm x}$ fuel, and this in turn is used to calculate the diffusion of oxygen within the fuel.

5. DEMONSTRATION PROBLEMS

A number of demonstration problems were developed to determine the degree of speedup provided by the re-initialization algorithm, as well as to ensure that use of this algorithm does not alter the calculated results.

5.1 Stand-Alone Thermochimica Calculations

The first test cases were for Thermochimica in stand-alone execution (i.e., no coupling). Two example systems were chosen for testing: a very simple carbon-oxygen system containing only C and O, and a much more complex nuclear fission product system including Pu, Np, U, Nd, Pr, Ce, La, Ba, Cs, Xe, I, Te, Pd, Rh, Ru, Tc, Mo, Zr, Y, Sr, Rb, O, and H. For each system timing was performed with and without enabling the re-initialization algorithm for three cases: repetition of an identical calculation; calculations with varying temperature; and calculations with varying element concentrations. The resulting speedups determined from each of these calculations are shown in Figure 2.

5.1.1 Carbon-Oxygen System

The carbon-oxygen example system contains 2 elements, 1 solution phase, and 4 pure phases. This is an extremely simple system, and therefore all Thermochimica calculations on this system are very brief. To produce a test approximately 1 minute in length, 2 million calls to Thermochimica must be made. This is naturally a highly unfavorable test case for the re-initialization algorithm, as a relatively large amount of calculation time is spent saving and loading data, and relatively little in the GEM solver loop. Therefore it is not surprising that small speedups or even slowdowns are provided by re-initialization in this case. This illustrates why re-initialization has been implemented as a setting available to the user rather than imposed

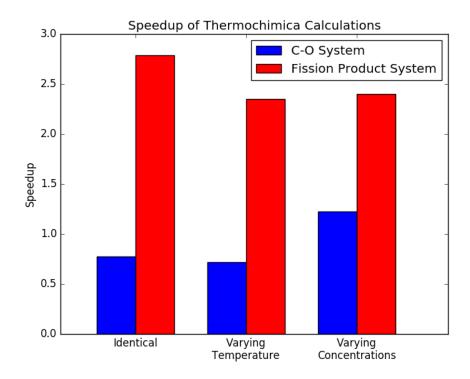


Figure 2. Speedup due to use of the re-initialization method for stand-alone Thermochimica calculations of the carbon-oxygen and fission product example systems.

automatically on all calculations: experienced users should be able to identify whether their combinations of system and available hardware makes re-initialization likely to be favorable, and use or omit it accordingly.

The base conditions for the carbon-oxygen example system were a temperature of 900 K, pressure of 1 atm, 1 mole of oxygen, and 2.4 moles of carbon. For the first test these conditions are unchanged between calculations. For the second case, the temperature was varied smoothly from 600 K to 1600 K over the 2 million calculations. In the third case, the amount of oxygen was varied smoothly from 1 mole to 6 moles over the calculation series. Surprisingly, only this last test shows speedup. Further testing revealed that this is because positive speedup is achieved for higher oxygen to carbon ratios. When there is more oxygen than carbon in the system then no pure condensed carbon phase is stable at 900 K, and the GEM solver takes a larger number of steps to converge. When the first test was run with 5 moles of carbon and all other conditions unchanged, the speedup was $1.16 \times$ rather than the $0.78 \times$ for the original test.

The lack of speedup – or even reduction in performance – in these cases can be understood when the number of GEM solver calls is considered. In the case of the identical calculation test, re-initialization results in exactly the same number of calls to the GEM solver as were made without re-initialization. Thus no speedup is provided by the algorithm, and the overhead of saving and loading the required re-initialization data results in an overall slowdown of the calculation.

5.1.2 Irradiated Nuclear Fuel

The second example system for stand-alone Thermochimica calculations made use of a database of uranium dioxide fuel with fission and activation products [1]. This database contains 23 elements, 15 solution phases, and 240 pure phases. Thermochimica calls for this system are much more expensive than for the carbon-oxygen system, and to keep the tests approximately the same length only 800 calculations were run per test for this example.

The base conditions for this example were a temperature of 900°C, pressure of 1 atm, and element concentrations as summarized in Table 2. These were again kept constant for the first test. For the second, the temperature was varied linearly from 600°C to 1400°C over the 800 calculations. In the third, the concentrations of all elements except U and O were varied linearly from 0 moles to 8× their base amounts over the calculation series. This very loosely approximates the evolution of fission products during burnup of uranium dioxide nuclear fuel. All three of these tests showed significant speedups, ranging from 2.62 to 2.97. For this system, the highest speedup was achieved for repetition of a calculation under constant conditions. This is the behaviour anticipated for the vast majority of cases: re-initialization ought to work best in cases where the conditions (and thus usually results) of a calculation are very similar to that of the preceding calculation. In contrast to the C-O system calculations, the number of calls to the GEM solver is greatly reduced when re-initialization is employed for this system. For the identical calculation example, the number of GEM solver calls is reduced by 76.2%, which implies that the GEM portion of the calculation is sped up approximately 4.2×, and the save/load overhead as well as non-accelerated portions of the calculations reduce the overall speedup to the 2.97× quoted above.

5.2 Bison Example Calculations

Previous coupling of Thermochimica to Bison has focused on calculations of the distribution and chemical potential of oxygen within the fluorite phase of uranium dioxide $(UO_{2\pm x})$ nuclear fuel [14, 13] for Light Water Reactor (LWR) applications. These example Bison calculations extend this work and demonstrate the reduction in simulation time for Bison objects that call Thermochimica when the re-initialization algorithm is employed.

These examples made use of a limited dataset containing only 3 chemical elements (O, U, Pu), 8 solution phases, and 18 stoichiometric phases, rather than that used in the above stand-alone example. Therefore based on the results of the stand-alone examples, we anticipate that greater speedup is possible for Bison simulations involving more chemically complex systems. The mobility model for non-stoichiometric uranium oxide from Moore *et al.* [6] and the thermodynamic treatment for the U-O system from Guéneau *et al.* [4] were used in these examples.

5.2.1 1D Bison Calculation

In the first Bison example, the diffusion of oxygen within uranium dioxide fuel with small amounts of plutonium is calculated on a 1-dimensional grid with 42 nodes. This system uses the U02XChemistryTransportSimplified AuxKernel and ThermochimicaNodalU02X UserObject described previously in Section 4. to call Thermochimica and extract important calculation results such as the concentration of interstitial oxygen and the chemical potential of oxygen. The calculation was run with

Table 2. Base concentrations of elements in fission product example system

Element	Moles
Pu	1×10^{-9}
Np	1×10^{-9}
U	5.6849×10^{-3}
Nd	1.8974×10^{-7}
Pr	1×10^{-9}
Ce	7.5956×10^{-8}
La	3.831×10^{-8}
Ba	9.9634×10^{-8}
Cs	1.2584×10^{-9}
Xe	1×10^{-9}
I	1×10^{-9}
Te	1×10^{-9}
Pd	1.0003×10^{-5}
Rh	7.3875×10^{-7}
Ru	4.2121×10^{-6}
Tc	1×10^{-9}
Mo	1.4263×10^{-5}
Zr	1.1667×10^{-7}
Y	1×10^{-9}
Sr	8.6762×10^{-7}
Rb	1×10^{-9}
O	1.1446707×10^{-2}
Н	1×10 ⁻⁹

and without re-initialization for 2.5 million seconds of simulation time (approximately one month) with time-steps of 10,000 seconds. The speedup achieved for this example was $2.72\times$.

5.2.2 3D Bison Calculation

A similar Bison simulation was performed on a 3-dimensional cylindrical mesh containing 2176 nodes. A parabolic, non-evolving temperature gradient was applied such that the cylinder was hottest in the centre (up to 2100 K), and cooler at the ends and edges (as low as 900 K) to induce redistribution of oxygen and gradients in the other Thermochimica output variables. This calculation was performed for 10 million seconds of simulation time with time-steps of 10,000 seconds on 16 cores on a shared memory workstation, with and without re-initialization enabled. The re-initialization algorithm achieved speedup of 7.38× for this example, shortening run-time from approximately 3 hours to as little as 25 minutes. This speedup is much greater than that measured for any of the previous examples, and was possible because the number of calls to the GEM solver routine was reduced by 96.1% when the re-initialization algorithm was employed. The speedup achieved for the entire calculation was much less than the implied 25× speedup of the GEM solver due to the overhead of saving and loading re-initialization data, as well as time spent in the remaining un-accelerated portions of the calculation.

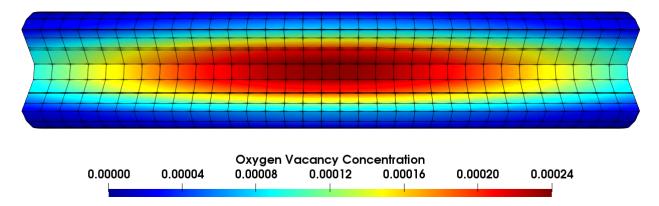


Figure 3. Cutaway view of oxygen vacancy concentration in uranium dioxide light water reactor fuel from a 3D Bison example calculation after 10 million seconds.

6. CONCLUSION

A method of re-initializing Thermochimica calculations within Bison based on the results of previous calculations has been developed. Subroutines have been provided to make use of this functionality both in the context of Thermochimica as a stand-alone software package and when Thermochimica is employed as a module in the MOOSE/Bison framework. We have developed a series of tests based on realistic nuclear fuel applications that demonstrate significant speedup of calculations due to the re-initialization algorithm, ranging from approximately 2.5× for most cases to as much as 7.38× for the largest test case: a 3-dimensional simulation in Bison of a uranium dioxide fuel element.

Along with these performance improvements, the generalization of the Thermochimica interface in Bison developed here will enable Bison developers to make use of Thermochimica in their calculations more efficiently and with less development overhead. ThermochimicaNodalData provides a common base class for all future development of MOOSE/Bison objects requiring Thermochimica calculations at nodal points in the FEM mesh. ThermochimicaNodalUO2X is an example derived UserObject that customizes the call to Thermochimica and stores Thermochimica output variables in a manner that allows access by Kernels and AuxKernels. UO2XChemistryTransportSimplified is an example of such and AuxKernel, and uses Thermochimica-calculated values such as the chemical potential of oxygen and the oxygen interstitial fraction to determine the rate of oxygen diffusion in uranium dioxide.

7. ACKNOWLEDGEMENTS

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